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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2008; month=1; day=11; hr=17; min=2; sec=10; ms=804;]

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Reviewer Comments:

Please realign amino acid number so that it lines up with every fifth
amino acid.

Application No: 10572522 Version No: 1.0

Input Set:

Output Set:

Started: 2007-12-18 14:57:25.234
Finished: 2007-12-18 14:57:26.071
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 837 ms
Total Warnings: 0
Total Errors: 5
No. of SeqIDs Defined: 1
Actual SeqID Count: 1

Error code	Error Description
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (20)
E 323	Invalid/missing amino acid numbering SEQID (1) POS (24)
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (25)
E 323	Invalid/missing amino acid numbering SEQID (1) POS (29)
E 323	Invalid/missing amino acid numbering SEQID (1)at Protein (30)

SEQUENCE LISTING

<110> Andreas BERGMANN

<120> METHOD FOR THE DIAGNOSIS OF DISEASES BY DETERMINING APOLIPOPROTEIN C-1

<130> 2582.011

<140> 10572522

<141> 2007-12-18

<150> PCT/EP2004/09965

<151> 2004-09-07

<150> DE 103 43 815.7

<151> 2003-09-22

<160> 1

<170> PatentIn version 2.1

<210> 1

<211> 57

<212> PRT

<213> Homo sapiens

<400> 1

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							20				25				30	

Glu	Leu	Ser	Ala	Lys	Met	Arg	Glu	Trp	Phe	Ser	Glu	Thr	Phe	Gln	Lys
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Val	Lys	Glu	Lys	Leu	Lys	Ile	Asp	Ser
				50			55	